

#7



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ENTERED

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/038,297

DATE: 12/26/2002

TIME: 15:14:34

Input Set : A:\10-038,297.asc.txt
Output Set: N:\CRF4\12262002\J038297.raw

1 <110> APPLICANT: Phan, Brigitte Chau
2 Virtanen, Jorma Antero
4 <120> TITLE OF INVENTION: Dual Bead Assays Including Covalent Linkages For
5 Improved Specificity And Related Optical Analysis Discs
7 <130> FILE REFERENCE: BTI2 00103501(USP2)US
9 <140> CURRENT APPLICATION NUMBER: 10/038,297
C--> 11 <141> CURRENT FILING DATE: 2002-12-10
W--> 13 <150> PRIOR APPLICATION NUMBER:
W--> 14 <151> PRIOR FILING DATE:
16 <160> NUMBER OF SEQ ID NOS: 2
18 <170> SOFTWARE: Microsoft Notepad Windows98
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 40
22 <212> TYPE: DNA
23 <213> ORGANISM: Artificial Sequence
W--> 25 <220> FEATURE: No Biological Significance.
W--> 26 Used for experimental purpose only.
W--> 27 <221> NAME/KEY:
28 <222> LOCATION:
29 <223> OTHER INFORMATION: Capture Probe
31 <400> SEQUENCE: 1
32 ccagtgaatt cgagctcggt acccgggat cctctagagt 40
34 <210> SEQ ID NO: 2
35 <211> LENGTH: 40
36 <212> TYPE: DNA
37 <213> ORGANISM: Artificial Sequence
W--> 39 <220> FEATURE: No Biological Significance.
W--> 40 Used for experimental purpose only.
W--> 41 <221> NAME/KEY:
42 <222> LOCATION:
43 <223> OTHER INFORMATION: Reporter Probe
45 <400> SEQUENCE: 2
46 cttagtcttt agatgcaagc ttggcgtaat catggtcata 40

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/038,297

DATE: 12/26/2002
TIME: 15:14:35

Input Set : A:\10-038,297.asc.txt
Output Set: N:\CRF4\12262002\J038297.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:13 M:256 W: Invalid Numeric Header Field, <150> PRIOR APPLICATION NUMBER:
L:14 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:25 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:26 M:259 W: Allowed number of lines exceeded, <220> FEATURE:
L:27 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:39 M:256 W: Invalid Numeric Header Field, <220> has non-blank data
L:40 M:259 W: Allowed number of lines exceeded, <220> FEATURE:
L:41 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2